### 中国横断山的地衣研究(6):灰梅属地衣修订\*

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摘要:通过对灰梅属(Canoparmelia)表型与基因型相结合的研究,澄清了云南南部报道的珊瑚大叶梅(Parmotrema conformatum)是裂芽灰梅(C. ecaperata)的错误鉴定。由于珊瑚大叶梅和裂芽灰梅在经典分类鉴别上容易混淆,通过 rDNA ITS 序列的系统发育分析,证明了裂芽灰梅与粉芽灰梅(C. texana)及同心灰梅(C. concrescens)有较近的亲缘关系;首次记录了粉芽灰梅在云南的分布范围;文中对两个新记录种进行了详细描述和讨论,并给出了中国已知灰梅属的检索表。

关键词: 灰梅属; 新记录; 地理分布; 分子系统学

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## Contributions to the Lichen Flora of the Hengduan Mountains, China (6): Revisional Study of the Genus *Canoparmelia* (Lichenized Ascomycota, Parmeliacaea)\*

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Abstract: A revision of Canoparmelia from the Hengduan Mountains (Yunnan, China) revealed the occurrence of C. ecaperata, based on specimens previously identified as Parmotrema conformatum. C. ecaperata is hereby reported as new to the flora of China. The study also led to the identification of C. texana as a component of the flora of Yunnan. In order to make sure the systematic position of these species, a phylogenetic analysis of rDNA ITS sequence was also carried out in this paper, indicating that C. ecaperata is closely related to C. texana and C. concrescens. Detailed information and discussion of these two new records are provided, together with a key to all the known taxa of Canoparmelia recorded from China. Following our revision, Parmotrema conformatum should be excluded from the flora of China.

Key words: Canoparmelia; New record; Geographic distribution; Molecular phylogeny

Canoparmelia Elix & Hale (Parmeliaceae, Lecanorales, Ascomycota) comprises ca. 40 species (Hawksworth et al., 1995; Thell et al., 2012) and is typified by C. texana (Tuck.) Elix & Hale. It was

segregated by Elix and Hale from *Pseudoparmelia* on the basis of morphological, geographic, ecological, and chemical characters (Elix *et al.*, 1986). Four species of *Canoparmelia* were previously known from

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China: two from Taiwan (Wei, 1991; Lai, 2000), and three from mainland China (Jia et al., 2008). As part of an inventory of the lichen flora of the Hengduan Mountains (Yunnan, China) several specimens of Canoparmelia were collected. These belong to C. ecaperata Elix & Hale, which is a new record to the flora of China, and to C. texana, which is new to Yunnan Province. All specimens of Parmotrema conformatum, previously reported from Yunnan were misidentified and belong to C. ecaperata.

### 1 Materials and Methods

The specimens were mainly collected from Yunnan Province, and were deposited in KUN (Herbarium of Kunming Institute of Botany). External morphological descriptions were based on air-dried materials, and observation under stereomicroscopy (Dongwon OSM-1). Chemical characteristics were examined by medullary color reaction, with the solution for the color reaction following the methods described by Yoshimura (1974): K (10% aqueous KOH solution), C (NaClO solution), KC (10% KOH followed by C solution), P (5% alcoholic P-phenylenediamine). Thin layer chromatography (TLC) of acetone extracts was also carried out to further characterize the chemical profile of the specimens, using solvent systems C and G (Culberson and Hör-Dur, 1970: Orange et al., 2001) and Lethariella cladonioides (Nyl.) Krog as control sample.

DNA was extracted from freshly collected and frozen herbarium specimens grinded to a fine powder, using Axyprep Multisource Genomic DNA Miniprep Kit following the manufacturer's instructions. The nuclear ITS (rDNA repeat) was amplified using primers ITS1 and ITS4 (White, 1990; Gardes and Bruns, 1993) in 25  $\mu$ L reaction volume composed of 2  $\mu$ L DNA, 1  $\mu$ L of each primer, 12.5  $\mu$ L 2×Taq PCR Mastermix (Aidlab) (contain Taq DNA Polymerase: 0.1 unit  $\cdot \mu$ L<sup>-1</sup>; MgCl<sub>2</sub>: 4 mmol  $\cdot$  L<sup>-1</sup>; dNTPs: 0.4 mmol  $\cdot$  L<sup>-1</sup>), 8.5  $\mu$ L dH<sub>2</sub>O. Amplifications were carried out in an automatic thermocycler (C 1000<sup>TM</sup>), with the following profile: initial denaturation at 94 °C

for 5 min, followed by 30 cycles, each of three steps (94  $^{\circ}$ C for 1 min, 56  $^{\circ}$ C for 1 min, 72  $^{\circ}$ C for 1.5 min), and a final extension at 72  $^{\circ}$ C for 7 min.

Twenty-seven sequences (18 retrieved from Gen-Bank and 9 obtained by authors, Table 1) were assembled into a matrix for phylogenetic analyses. The sequence database was subjected to automated alignment and then manually inspected in BioEdit. Hypotrachynia adducta was set as the outgroup. Ambiguous regions were deleted. Bayesian inference of phylogeny was performed with in MrBayes v3. 1. 2. ModelTest 3.7 was used to estimate separate best-fit models of evolution for ITS. Bayesian analyses were carried out using the Metropoliscoupled Markov chain Monte Carlo method (MCMCMC). Analyses were run under the GTR model using a gamma-distributed rate parameter and a proportion of invariable sites. Two parallel MCMCMC runs were performed, each using four chains and 1 million generations, sampling trees of every 100th generation. The initial burn-in was set to 50%. A 50% majority-rule consensus cladogram was computed from the remaining trees; the proportions of this tree correspond to Bayesian posterior probabilities (BPP), and clades with PP≥0.95 were considered to be significantly supported. The phylogenic trees are shown in Fig. 1.

### 2 Result

# 2. 1 Key to the species of known *Canoparmelia* in China

····· C. ecaperata

Table 1 Species and specimens of Parmeliaceae analyzed

Species name	Accession NO.	Location	Source	Species name	Accession NO.	Location	Source
Canoparmelia concrescens	GU994543	Kenya	GenBank	F. papillosa	HM010936	Uruguay	GenBank
C. ecaperata	KC978843	China	10-31934	F. soredians	AY586562	Spain	GenBank
C. ecaperata	KC978844	China	12-33440	F. springtonensis	EF042907	Australia	GenBank
C. ecaperata	KC978845	China	12-33611	F. subambigua	GU994551	Argentina	GenBank
C. ecaperata	KC978846	China	12-34474	F. subambigua	JN943839	Argentina	GenBank
C. nairobiensis	GU994545	Kenya	GenBank	Hypotrachynia adducta	KC978855	China	12-34288
C. texana	KC978847	China	12-37655	Parmotrema subtinctorium	KC978853	China	12-35202
C. texana	KC978848	China	12-37665	P. norsticticatum	GU994577	South Africa	GenBank
C. texana	GU994547	Kenya	GenBank	P. reticulatum	AY586579	Portugal	GenBank
Flavoparmelia baltimorensis	AY586559	USA	GenBank	P. subcaperatum	AY586557	Australia	GenBank
F. caperata	AY581059	Spain	GenBank	P. subtinctorium	GU593037	China	GenBank
F. citrinescens	GU994550	Argentina	GenBank	P. tinctorum	AY586570	Australia	GenBank
F. haywardiana	HM010933	New Zealand	GenBank	P. tinctorun	KC978854	China	12-33444
F. marchantii	JN943840		GenBank				

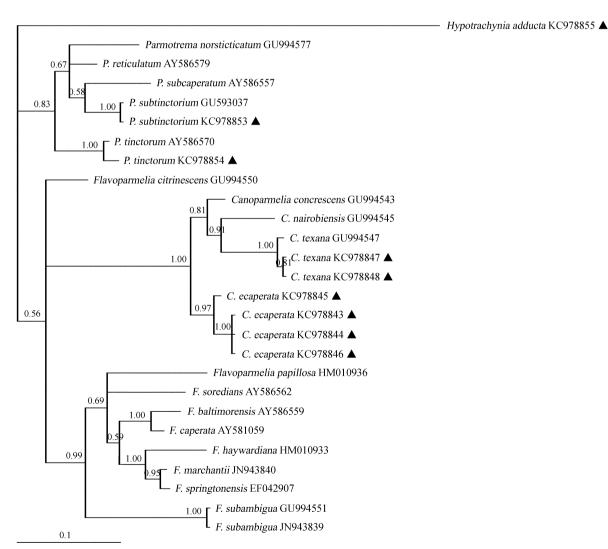


Fig. 1 Phylogenic construction of the systematic position of C. ecaperata based on rDNA sequence date.  $\blacktriangle$  Sequences obtained by author

### 2. 2 Taxonomy

Canoparmelia ecaperata Elix & Hale in Mycotaxon 27: 277, 1986. ——Parmelia ecaperata Müll. Arg. in Flora, Jena 74: 378. 1891. ——Pseudoparmelia ecaperata (Müll. Arg.) Hale in Phytologia 29: 190, 1974.

**Diagnostic characters:** Thallus foliose, adnate on bark, grayish yellow to yellow-green,  $4-14\,\mathrm{cm}$  in diam.; lobes irregularly sublinear,  $1-3\,\mathrm{mm}$  wide, without cilia; upper surface plane, continuous, sometimes cracked on the old part; covered with dense isidia, simple and short,  $0.1-1\,\mathrm{mm}$  long, cylindrical or clavate, mature ones with brown to black tips; medulla white,  $62.5-75\,\mathrm{\mu m}$  thick; lower surface

black, turning brownish near the margin, rhizines rare, simple and black, 0.5-1.5 mm long; lower cortex 25-75 µm thick, Apothecia not seen (Fig. 2).

**Chemistry:** Cortex K-, medulla K-, C- (pink), KC-, P-; atranorin, divaricatic acid, usnic acid and an unknown compound (Rf value between classes 4 and 5).

**Distribution:** China: Yunnan Province, Nanjian Co. (Fig. 3); Africa, Nepal, India, Thailand (Hale and Mason, 1976; Singh and Sinha, 2010; Swinscow *et al.*, 1988; Wolseley *et al.*, 2002).

**Habitat and ecology:** This species grows on the bark of *Pinus yunnanensis* from 1 300 m to 2 100 m in Nanjian Co., Yunnan.

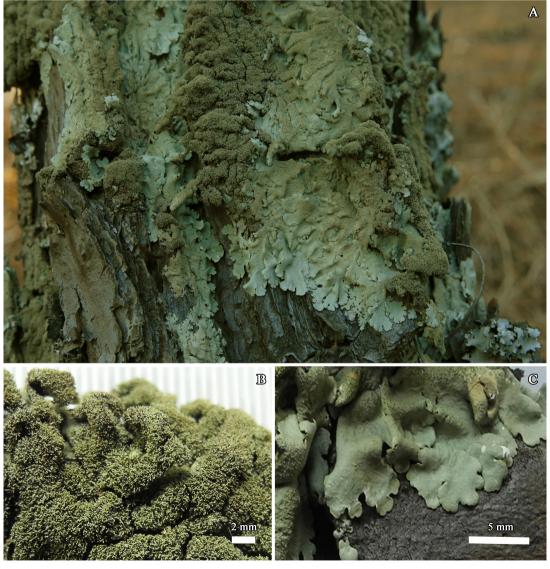


Fig. 2 Canoparmelia ecaperata. A. Habitat; B. Cylindrical laminal isidia. C. Narrow and sublinear lobes. (All based on Li S. Wang 12-37823)

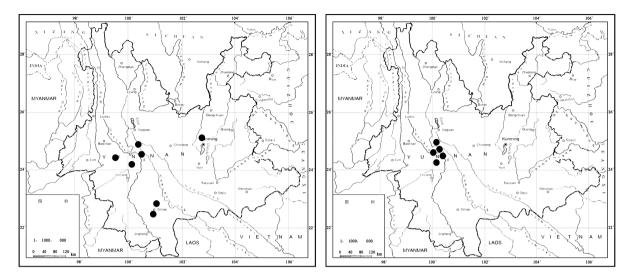


Fig. 3 Distribution map of C. ecaperata (left) and C. texana (right) from Hengduan Mountains

**Remarks**: Canoparmelia ecaperata is similar to Parmotrema conformatum and Flavoparmelia caperata. The former differs from C. ecaperata in the presence of fumarprotocetraric and protocetraric acids (P+ red), ciliate lobes (Chen et al., 2005; Benatti and Marcelli, 2009). Moreover, C. ecaperata occurs mainly in the tropical area of the Southern Hemisphere (Elix and McCarthy, 1998; Nöske and Sipman, 2004), whereas C. ecaperata has been reported from India and Thailand (Hale, 1976). Flavoparmelia caperata differs in the sorediate rather than isidiate lobes, and the presence of protocetraric acid rather than atranorin as the main compound. Canoparmelia concrescens might be confused with C. ecaperata but is distinguished by the bright yellow thallus (containing usnic acid in the cortex), and its distribution only in Africa. Finally C. amazonica, which also has isidia can be easily distinguished by the lack of usnic acid and spare rhizines.

**Specimens examined: Yunnan Province:** Nanjian Co.: Leqiu Village, N: 25°00′25.29″, E: 100° 21′43.50″, alt. 2 060 m, on *Pinus yunnanensis* bark, Wang Lisong & Wang Xinyu 12–37814, 12–37820, 12–37822, 12–37823, 21 Dec. 2012; Gonglang Village, N: 24°43.103′, E: 100°25.514′, alt. 1 840 m, on *Pinus yunnanensis* bark, Wang Lisong *et al.* 12– 33611, 25 March 2012. Longling Co., Songshan Mt., N: 24°44.672′, E: 98°54.281′, alt. 2 000 m, on *Pinus*, Wang Lisong *et al.* 12-33440, 30 March 2012; Langcang Co., Mengla Village, on *Pinus*, alt. 1 100 m, Wang Lisong 10-31934, 11 Dec. 2010; Pu'er City, Ning'er Co., nearby road 213, N: 22°53′55. 44″, E: 101°02′28. 41″, alt. 1 334 m, on bark, Wang Lisong *et al.* 12-34474, 21 June 2012; Pu'er City, N: 22. 48°, E: 100. 58°, Zhao Ji-ding & Chen Yu-ben 2942, HMAS 100308, 18 Nov. 1960; Kunming City, Heilongtan, N: 25. 04°, E: 102. 42°, Zhao Ji-ding & Chen Yu-ben 4748, HMAS 100307, 16 Dec. 1960.

Canoparmelia texana (Tuck.) Elix & Hale in Mycotaxon 27: 278, 1986. ——Parmelia texana Tuckerman in American Journal of Science and Arts series 2 (253): 424, 1858. ——Pseudoparmelia texana (Tuckerman) Hale in Phytologia 28: 191, 1974.

**Diagnostic characters:** Thallus gray, upper surface covered with dense soralia, punctiform or capitate; lower surface black; moderately rhizinate, rhizines simple; containing atranorin and divaricatic acid.

**Distribution:** Pantemperate outside of Europe and pantropical at higher elevations. Shandong and Taiwan Province in China (Jia, 2008; Lai, 2000); New to Yunnan Province (Fig. 3).

**Specimens examined: Yunnan Province**, Nanjian Co.: Leqiu Village. N: 25°00′25. 29″, E: 100° 21′43. 50″, alt. 2 060 m, on *Pinus yunnanensis* bark,

Wang Lisong & Wang Xinyu 12-37805, 12-37810, 12-37889, 21 Dec. 2012; Wuliang Medicine Valley, N: 24°52′2.76″; E: 100°34′51.39″, alt. 2 345 m, on Pinus amandii bark, Wang Lisong & Wang Xinyu 12-37655, 19 Dec. 2012; Hongxing Village, Wuliang Medicine Valley, N: 24°50′40.41″, E: 100° 38'4.52", alt. 2 341 m, on bark, Wang Lisong & Wang Xinyu 12 - 36923, 16 Oct. 2012; Gonglang Country, Jinguo Village, N: 24°48.557′, E: 100° 24. 839', alt. 2 270 m, on bark, Wang Lisong et al. 12-33742, 25 March 2012; Ziqiang Village, N: 24°46′10.55″; E: 100°25′08.13″, alt. 1 640 m, on bark, Wang Lisong & Wang Xinyu 12-37674, 19 Dec. 2012; Ziqiang Village, N: 24°47′21.60″; E: 100°25′07.78″, alt. 1 740 m, on bark, Wang Lisong & Wang Xinyu 12-37665, 19 Dec. 2012; Dadianshan Mt., N: 24°51′48″, E: 100°34′39″, alt. 2543 m, on branch, Wang Lisong et al. 12-32957, 21 March 2012.

Canoparmelia amazonica (Nyl.) Elix & Hale in Mycotaxon 27: 278, 1986

Thallus grey, upper surface covered with rare branched isidia, rhizines dense. Containing atranorin and protocetraric acid (Hale, 1976).

Canoparmelia crozalsiana (Bouly de Lesdain) Elix & Hale in Mycotaxon 27: 278, 1986

Thallus greenish or gray, sorediate, moderately rhizinate. Containing atranorin, stictic acid, constictic acid (Hale, 1976).

Canoparmelia subtiliacea (Nyl.) Elix & Hale in Mycotaxon 27: 279, 1986

Thallus grey, upper surface without soredia or isidia, rhizines rare. Containing atranorin and caperatic acid (Hale, 1976).

### 3 Discussion

Since the genera Flavoparmelia, Pamotrema, and Canoparmelia are similar in external morphology (Table 2) and therefore hard to distinguish from each other, the phylogenetic method was applied in this study to verify the systematic position of these taxa and validate the morphological identification. The genus Parmotrema is characterized by broad lobes (usually more than 5 mm) and rare, simple rhizines in the central part of the lower surface, upper surface grayish green, without usnic acid (Chen et al., 2005); Flavoparmelia is characterized by broad and roundish lobes, containing usnic acid on the upper surface so that the color is yellow-green (Hale, 1986): Canoparmelia has sublinear and rather narrow lobes, closely attached to the substrate, has a subtropical to tropical distribution, and always contains atranorin as the main compound (Elix et al., 1986).

Parmotrema conformatum was previously reported from Yunnan Province by Zhao et al. (1982), based, however, on two specimens (HMAS 100307 100308) that lacked cilia and had P- medullary reaction. Careful examination of the two specimens revealed that they belong to C. ecaperata according to morphological, chemical, and distributional characters. This is further supported by inferences from molecular data whereby both specimens belong to Canoparmelia and to C. ecaperata in particular.

Table 2 Key differences among Can	parmelia, Flavoparmelia and Parmotrema
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	Flavo parmelia	Parmotrema	Can opar melia
Thallus color	Yellowish-green	Grey to greenish	Grey to yellowish
Lobes	Broad (2-6 mm)	Rather broad (5-10 mm)	Medium (0.5-5 mm)
Lower surface	Black, with a narrow, brown, naked marginal zone	Brown to black, with broad naked zone.	Black, narrow, brown, marginal zone ( $< 1 \text{ mm}$ )
Rhizines	Sparse to moderate, simple or branched	Sparse and simple, in the thallus center	Moderate to dense, Simple or tufted
Characteristic compounds	Usnic acid	Atranorin	Atranorin, usnic acid
Distribution	Temperate	Temperate to tropical	Subtropical to tropical

The phylogenetic tree showed a final alignment consisting of 480 characters, of which 281 characters are constant, 31 variable parsimony-uninformative, and 168 parsimony-informative. All characters have equal weight. Gaps are treated as missing. The GC content of *C. ecaperata* is 54.7%, lower than *C. texana* (56.2%). The molecular data showed that the phylogenetic relationship of *C. ecaperata* is closer to *Canoparmelia*, which is in accord with our morphological identification. The final result confirms the placement of this species in the *Canoparmelia* clade. In this analysis, *C. ecaperata* is recovered as sister group of *C. texana* and *C. concrescens* with high support.

In conclusion, the presence of two taxa of *Canoparmelia* has been confirmed from the Hengduan Mountains based on morphological, chemical and phylogenetic study: *C. ecaperata* is reported as new to China and *C. texana* as new to Yunnan. The previous record of *Parmotrema conformatum* from China is based on misidentification (= *C. ecaperata*) which implies that the species should be deleted from the flora of China. The study added to the lichen biodiversity of the Hengduan Mountains, and also showed that there is still a high potential for new discoveries in this region.

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